

[illegible]

1.1.8	77.9	3815	6	US-09-724-676-4798	Ap	Sequence 4798	c 607	1.1.8	77.9	3815	1	US-10-080-414-1414-1138	Ap	Sequence 4141	c 607
1.1.8	77.9	3816	6	US-09-724-676-4798	Ap	Sequence 4798	c 608	1.1.8	77.9	3816	1	US-10-080-414-1414-1139	Ap	Sequence 4141	c 608
1.1.8	77.9	3817	6	US-09-724-676-4777	Ap	Sequence 4777	c 609	1.1.8	77.9	3817	1	US-10-080-414-1414-1140	Ap	Sequence 4141	c 609
1.1.8	77.9	3818	6	US-09-724-676-4777	Ap	Sequence 4777	c 610	1.1.8	77.9	3818	1	US-10-080-414-1414-1141	Ap	Sequence 4141	c 610
1.1.8	77.9	3819	6	US-09-724-676-4777	Ap	Sequence 4777	c 611	1.1.8	77.9	3819	1	US-10-080-414-1414-1142	Ap	Sequence 4141	c 611
1.1.8	77.9	3820	6	US-09-724-676-4777	Ap	Sequence 4777	c 612	1.1.8	77.9	3820	1	US-10-080-414-1414-1143	Ap	Sequence 4141	c 612
1.1.8	77.9	3821	6	US-09-724-676-4777	Ap	Sequence 4777	c 613	1.1.8	77.9	3821	1	US-10-080-414-1414-1144	Ap	Sequence 4141	c 613
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1.1.8	77.9	3823	6	US-09-724-676-4777	Ap	Sequence 4777	c 615	1.1.8	77.9	3823	1	US-10-080-414-1414-1146	Ap	Sequence 4141	c 615
1.1.8	77.9	3824	6	US-09-724-676-4777	Ap	Sequence 4777	c 616	1.1.8	77.9	3824	1	US-10-080-414-1414-1147	Ap	Sequence 4141	c 616
1.1.8	77.9	3825	6	US-09-724-676-4777	Ap	Sequence 4777	c 617	1.1.8	77.9	3825	1	US-10-080-414-1414-1148	Ap	Sequence 4141	c 617
1.1.8	77.9	3826	6	US-09-724-676-4777	Ap	Sequence 4777	c 618	1.1.8	77.9	3826	1	US-10-080-414-1414-1149	Ap	Sequence 4141	c 618
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1.1.8	77.9	3829	6	US-09-724-676-4777	Ap	Sequence 4777	c 621	1.1.8	77.9	3829	1	US-10-080-414-1414-1152	Ap	Sequence 4141	c 621
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1.1.8	77.9	3833	6	US-09-724-676-4777	Ap	Sequence 4777	c 625	1.1.8	77.9	3833	1	US-10-080-414-1414-1156	Ap	Sequence 4141	c 625
1.1.8	77.9	3834	6	US-09-724-676-4777	Ap	Sequence 4777	c 626	1.1.8	77.9	3834	1	US-10-080-414-1414-1157	Ap	Sequence 4141	c 626
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1.1.8	77.9	3838	6	US-09-724-676-4777	Ap	Sequence 4777	c 630	1.1.8	77.9	3838	1	US-10-080-414-1414-1161	Ap	Sequence 4141	c 630
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1.1.8	77.9	3840	6	US-09-724-676-4777	Ap	Sequence 4777	c 632	1.1.8	77.9	3840	1	US-10-080-414-1414-1163	Ap	Sequence 4141	c 632
1.1.8	77.9	3841	6	US-09-724-676-4777	Ap	Sequence 4777	c 633	1.1.8	77.9	3841	1	US-10-080-414-1414-1164	Ap	Sequence 4141	c 633
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1.1.8	77.9	3849	6	US-09-724-676-4777	Ap	Sequence 4777	c 641	1.1.8	77.9	3849	1	US-10-080-414-1414-1172	Ap	Sequence 4141	c 641
1.1.8	77.9	3850	6	US-09-724-676-4777	Ap	Sequence 4777	c 642	1.1.8	77.9	3850	1	US-10-080-414-1414-1173	Ap	Sequence 4141	c 642
1.1.8	77.9	3851	6	US-09-724-676-4777	Ap	Sequence 4777	c 643	1.1.8	77.9	3851	1	US-10-080-414-1414-1174	Ap	Sequence 4141	c 643
1.1.8	77.9	3852	6	US-09-724-676-4777	Ap	Sequence 4777	c 644	1.1.8	77.9	3852	1	US-10-080-414-1414-1175	Ap	Sequence 4141	c 644
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1.1.8	77.9	3856	6	US-09-724-676-4777	Ap	Sequence 4777	c 648	1.1.8	77.9	3856	1	US-10-080-414-1414-1179	Ap	Sequence 4141	c 648
1.1.8	77.9	3857	6	US-09-724-676-4777	Ap	Sequence 4777	c 649	1.1.8	77.9	3857	1	US-10-080-414-1414-1180	Ap	Sequence 4141	c 649
1.1.8	77.9	3858	6	US-09-724-676-4777	Ap	Sequence 4777	c 650	1.1.8	77.9	3858	1	US-10-080-414-1414-1181	Ap	Sequence 4141	c 650
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1.1.8	77.9	3865	6	US-09-724-676-4777	Ap	Sequence 4777	c 657	1.1.8	77.9	3865	1	US-10-080-414-1414-1188	Ap	Sequence 4141	c 657
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1.1.8	77.9	3871	6	US-09-724-676-4777	Ap	Sequence 4777	c 663	1.1.8	77.9	3871	1	US-10-080-414-1414-1194	Ap	Sequence 4141	c 663
1.1.8	77.9	3872	6	US-09-724-676-4777	Ap	Sequence 4777	c 664	1.1.8	77.9	3872	1	US-10-080-414-1414-1195	Ap	Sequence 4141	c 664
1.1.8	77.9	3873	6	US-09-724-676-4777	Ap	Sequence 4777	c 665	1.1.8	77.9	3873	1	US-10-080-414-1414-1196	Ap	Sequence 4141	c 665
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1.1.8	77.9	3876	6	US-09-724-676-4777	Ap	Sequence 4777	c 668	1.1.8	77.9	3876	1	US-10-080-414-1414-1199	Ap	Sequence 4141	c 668
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1.1.8	77.9	3878	6	US-09-724-676-4777	Ap	Sequence 4777	c 670	1.1.8	77.9	3878	1	US-10-080-414-1414-1201	Ap	Sequence 4141	c 670
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1.1.8	77.9	3880	6	US-09-724-676-4777	Ap	Sequence 4777	c 672	1.1.8	77.9	3880	1	US-10-080-414-1414-1203	Ap	Sequence 4141	c 672
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1.1.8	77.9	3883	6	US-09-724-676-4777	Ap	Sequence 4777	c 675	1.1.8	77.9	3883	1	US-10-080-414-1414-1206	Ap	Sequence 4141	c 675
1.1.8	77.9	3884	6	US-09-724-676-4777	Ap	Sequence 4777	c 676	1.1.8	77.9	3884	1	US-10-080-414-1414-1207	Ap	Sequence 4141	c 676
1.1.8	77.9	3885	6	US-09-724-676-4777	Ap	Sequence 4777	c 677	1.1.8	77.9	3885	1	US-10-080-414-1414-1208	Ap	Sequence 4141	c 677
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[illegible]


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1  CURRENT APPLICATION NUMBER: 05/06/24.676
2  CURRENT FILING DATE: 2000.11.28
3  NUMBER OF SEQ ID NOS: 97222
4  SOFTWARE: Patent In Version 3.2
5  SEQ ID NO: 1454
6  LENGTH: 776
7  TYPE: DNA
8  ORGANISM: Homo sapiens
9  1 1454 1454

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Query Match: 100.0% Score: 19.18 E-04
Best Local Similarity: 100.0% Prod. No. 45
Matches: 19 Conserved: 19 Mismatches: 0

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14 1 1454AAGAGATGAAAGG 19

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1  CURRENT APPLICATION NUMBER: 05/06/24.676
2  CURRENT FILING DATE: 2000.11.28
3  NUMBER OF SEQ ID NOS: 97222
4  SOFTWARE: Patent In Version 3.2
5  SEQ ID NO: 1454
6  LENGTH: 776
7  TYPE: DNA
8  ORGANISM: Homo sapiens
9  1 1454 1454

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Query Match: 100.0% Score: 19.18 E-04
Best Local Similarity: 100.0% Prod. No. 45
Matches: 19 Conserved: 19 Mismatches: 0

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14 1 1454AAGAGATGAAAGG 19

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1  CURRENT APPLICATION NUMBER: 05/06/24.676
2  CURRENT FILING DATE: 2000.11.28
3  NUMBER OF SEQ ID NOS: 97222
4  SOFTWARE: Patent In Version 3.2
5  SEQ ID NO: 1454
6  LENGTH: 776
7  TYPE: DNA
8  ORGANISM: Homo sapiens
9  1 1454 1454

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Query Match: 100.0% Score: 19.18 E-04
Best Local Similarity: 100.0% Prod. No. 45
Matches: 19 Conserved: 19 Mismatches: 0

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14 1 1454AAGAGATGAAAGG 19

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1  CURRENT APPLICATION NUMBER: 05/06/24.676
2  CURRENT FILING DATE: 2000.11.28
3  NUMBER OF SEQ ID NOS: 97222
4  SOFTWARE: Patent In Version 3.2
5  SEQ ID NO: 1454
6  LENGTH: 776
7  TYPE: DNA
8  ORGANISM: Homo sapiens
9  1 1454 1454

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Query Match: 100.0% Score: 19.18 E-04
Best Local Similarity: 100.0% Prod. No. 45
Matches: 19 Conserved: 19 Mismatches: 0

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1  CURRENT APPLICATION NUMBER: 05/06/24.676
2  CURRENT FILING DATE: 2000.11.28
3  NUMBER OF SEQ ID NOS: 97222
4  SOFTWARE: Patent In Version 3.2
5  SEQ ID NO: 1454
6  LENGTH: 776
7  TYPE: DNA
8  ORGANISM: Homo sapiens
9  1 1454 1454

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Query Match: 100.0% Score: 19.18 E-04
Best Local Similarity: 100.0% Prod. No. 45
Matches: 19 Conserved: 19 Mismatches: 0

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27 1 1454AAGAGATGAAAGG 19

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14 1 1454AAGAGATGAAAGG 19

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1  CURRENT APPLICATION NUMBER: 05/06/24.676
2  CURRENT FILING DATE: 2000.11.28
3  NUMBER OF SEQ ID NOS: 97222
4  SOFTWARE: Patent In Version 3.2
5  SEQ ID NO: 1454
6  LENGTH: 776
7  TYPE: DNA
8  ORGANISM: Homo sapiens
9  1 1454 1454

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Matches: 19 Conserved: 19 Mismatches: 0

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1  CURRENT APPLICATION NUMBER: 05/06/24.676
2  CURRENT FILING DATE: 2000.11.28
3  NUMBER OF SEQ ID NOS: 97222
4  SOFTWARE: Patent In Version 3.2
5  SEQ ID NO: 1454
6  LENGTH: 776
7  TYPE: DNA
8  ORGANISM: Homo sapiens
9  1 1454 1454

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Best Local Similarity: 100.0% Prod. No. 45
Matches: 19 Conserved: 19 Mismatches: 0

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27 1 1454AAGAGATGAAAGG 19

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14 1 1454AAGAGATGAAAGG 19

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Sun May 25 14:51:40 2003

11 06 03AAGAGAGATGAAAGG 77

RESULT 35

US-09-724-676-14548

Sequence 14548, Application US/09724676

GENERAL INFORMATION:

APPLICANT: Compugen LTD

TITLE OF INVENTION: Variants of alternative splicing

FILE REFERENCE: 129181.4 Compugen

CURRENT APPLICATION NUMBER: us/09/724-676

CURRENT FILING DATE: 2000-11-28

NUMBER OF SEQ ID NOS: 97222

SOFTWARE: Patent in version 3.2

SEQ ID NO: 14548

LENGTH: 815

TYPE: DNA

ORGANISM: Homo sapiens

US-09-724-676-14548

Query Match

Best Local Similarity 100.0%; Score 19; DB 6; Length 815;

Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

09 05 03AAGAGAGATGAAAGG 77

RESULT 41

US-09-724-676-14548

Sequence 14548, Application US/09724676

GENERAL INFORMATION:

APPLICANT: Compugen LTD

TITLE OF INVENTION: Variants of alternative splicing

FILE REFERENCE: 129181.4 Compugen

CURRENT APPLICATION NUMBER: us/09/724-676

CURRENT FILING DATE: 2000-11-28

NUMBER OF SEQ ID NOS: 97222

SOFTWARE: Patent in version 3.2

SEQ ID NO: 14548

LENGTH: 815

TYPE: DNA

ORGANISM: Homo sapiens

US-09-724-676-14548

Query Match

Best Local Similarity 100.0%; Score 19; DB 6; Length 815;

Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

09 05 03AAGAGAGATGAAAGG 77

RESULT 35

US-09-724-676-14555

Sequence 14555, Application US/09724676

GENERAL INFORMATION:

APPLICANT: Compugen LTD

TITLE OF INVENTION: Variants of alternative splicing

FILE REFERENCE: 129181.4 Compugen

CURRENT APPLICATION NUMBER: us/09/724-676

CURRENT FILING DATE: 2000-11-28

NUMBER OF SEQ ID NOS: 97222

SOFTWARE: Patent in version 3.2

SEQ ID NO: 14555

LENGTH: 844

TYPE: DNA

ORGANISM: Homo sapiens

US-09-724-676-14555

Query Match

Best Local Similarity 100.0%; Score 19; DB 6; Length 844;

Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Best Local Similarity 100.0%; Score 19; DB 6; Length 844;

Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

09 05 03AAGAGAGATGAAAGG 77

10 05 03AAGAGAGATGAAAGG 77

RESULT 41

US-09-724-676-14555

Sequence 14555, Application US/09724676

GENERAL INFORMATION:

APPLICANT: Compugen LTD

TITLE OF INVENTION: Variants of alternative splicing

FILE REFERENCE: 129181.4 Compugen

CURRENT APPLICATION NUMBER: us/09/724-676

CURRENT FILING DATE: 2000-11-28

NUMBER OF SEQ ID NOS: 97222

SOFTWARE: Patent in version 3.2

SEQ ID NO: 14555

LENGTH: 844

TYPE: DNA

ORGANISM: Homo sapiens

US-09-724-676-14555

Query Match

Best Local Similarity 100.0%; Score 19; DB 6; Length 844;

Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

09 05 03AAGAGAGATGAAAGG 77

10 05 03AAGAGAGATGAAAGG 77

RESULT 41

US-09-724-676-14555

Sequence 14555, Application US/09724676

GENERAL INFORMATION:

APPLICANT: Compugen LTD

TITLE OF INVENTION: Variants of alternative splicing

FILE REFERENCE: 129181.4 Compugen

CURRENT APPLICATION NUMBER: us/09/724-676

CURRENT FILING DATE: 2000-11-28

NUMBER OF SEQ ID NOS: 97222

SOFTWARE: Patent in version 3.2

SEQ ID NO: 14555

LENGTH: 844

TYPE: DNA

ORGANISM: Homo sapiens

US-09-724-676-14555

Query Match

Best Local Similarity 100.0%; Score 19; DB 6; Length 844;

Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

09 05 03AAGAGAGATGAAAGG 77

10 05 03AAGAGAGATGAAAGG 77

RESULT 45

US-09-724-676-14555

Sequence 14555, Application US/09724676

GENERAL INFORMATION:

APPLICANT: Compugen LTD

TITLE OF INVENTION: Variants of alternative splicing

FILE REFERENCE: 129181.4 Compugen

CURRENT APPLICATION NUMBER: us/09/724-676

CURRENT FILING DATE: 2000-11-28

NUMBER OF SEQ ID NOS: 97222

SOFTWARE: Patent in version 3.2

SEQ ID NO: 14555

LENGTH: 844

GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676A
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: Patent in version 3.2
SEQ ID NO 14542
LENGTH: 904
TYPE: DNA
ORGANISM: Homo sapiens
US-09-724-676A-14542

Query Match 100.0% Score 19; DB 6; Length 904;
Best Local Similarity 100.0%; Pred. No. 45;
Matches 19; Conservative 0; Mismatches 0; Gaps 0;
27 1 GGAAGAGAGATGAGACCC 19
28 5 GGAAGAGAGATGAGACCC 77

RESULT 42
US-09-724-676-14552/C
Sequence 14552, Application US/09/724676
GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: Patent in version 3.2
SEQ ID NO 14552
LENGTH: 918
TYPE: DNA
ORGANISM: Homo sapiens
US-09-724-676-14552

Query Match 100.0% Score 19; DB 6; Length 918;
Best Local Similarity 100.0%; Pred. No. 45;
Matches 19; Conservative 0; Mismatches 0; Gaps 0;
27 1 GGAAGAGAGATGAGACCC 19
28 5 GGAAGAGAGATGAGACCC 77

RESULT 43
US-09-724-676A-14552/C
Sequence 14552, Application US/09/724676A
GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676A
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: Patent in version 3.2
SEQ ID NO 14552
LENGTH: 918
TYPE: DNA
ORGANISM: Homo sapiens
US-09-724-676A-14552

Query Match 100.0% Score 19; DB 6; Length 918;
Best Local Similarity 100.0%; Pred. No. 45;
Matches 19; Conservative 0; Mismatches 0; Gaps 0;
27 1 GGAAGAGAGATGAGACCC 19
28 5 GGAAGAGAGATGAGACCC 77

RESULT 44
US-09-724-676-14518/C
Sequence 14518, Application US/09/724676
GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: Patent in version 3.2
SEQ ID NO 14518
LENGTH: 927
TYPE: DNA
ORGANISM: Homo sapiens
US-09-724-676-14518

Query Match 100.0% Score 19; DB 6; Length 927;
Best Local Similarity 100.0%; Pred. No. 45;
Matches 19; Conservative 0; Mismatches 0; Gaps 0;
27 1 GGAAGAGAGATGAGACCC 19
28 5 GGAAGAGAGATGAGACCC 77

RESULT 45
US-09-724-676A-14518/C
Sequence 14518, Application US/09/724676A
GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676A
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: Patent in version 3.2
SEQ ID NO 14518
LENGTH: 927
TYPE: DNA
ORGANISM: Homo sapiens
US-09-724-676A-14518

Query Match 100.0% Score 19; DB 6; Length 927;
Best Local Similarity 100.0%; Pred. No. 45;
Matches 19; Conservative 0; Mismatches 0; Gaps 0;
27 1 GGAAGAGAGATGAGACCC 19
28 5 GGAAGAGAGATGAGACCC 77

RESULT 46
US-09-724-676-14517/C
Sequence 14517, Application US/09/724676
GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: Patent in version 3.2
SEQ ID NO 14517
LENGTH: 934
TYPE: DNA
ORGANISM: Homo sapiens
US-09-724-676-14517

Query Match 100.0% Score 19; DB 6; Length 934;
Best Local Similarity 100.0%; Pred. No. 45;
Matches 19; Conservative 0; Mismatches 0; Gaps 0;
27 1 GGAAGAGAGATGAGACCC 19
28 5 GGAAGAGAGATGAGACCC 77

Matched 100 Conserved 100 Mismatches 00 Length 00

Query Match
Best Local Similarity 100.00%
Method 100.00% 100.00%

Result 10

us-09-925-548-1454/0
1 Response 145.0% Application US/09/240/6A
2 GENERAL INFORMATION:
3 APPLICANT: Compugen LTD
4 TITLE OF INVENTION: Variants of alternative splicing
5 FILE REFERENCE: 129181.4 Compugen
6 CURRENT APPLICATION NUMBER: US/09/240/6A
7 NUMBER OF SEQ ID NOS: 97222
8 SOFTWARE: Patent in version 3.2
9 SEQ ID NO: 1454
10 LENGTH: 964
11 TYPE: DNA
12 ORGANISM: Homo sapiens
US-09-925-548-1454

Query Match
Best Local Similarity 100.00%
Method 100.00% 100.00%

Query Match
Best Local Similarity 100.00%
Method 100.00% 100.00%

Result 10

us-09-925-548-1454/0
1 Response 145.0% Application US/09/240/6A
2 GENERAL INFORMATION:
3 APPLICANT: Compugen LTD
4 TITLE OF INVENTION: Variants of alternative splicing
5 FILE REFERENCE: 129181.4 Compugen
6 CURRENT APPLICATION NUMBER: US/09/240/6A
7 NUMBER OF SEQ ID NOS: 97222
8 SOFTWARE: Patent in version 3.2
9 SEQ ID NO: 1454
10 LENGTH: 964
11 TYPE: DNA
12 ORGANISM: Homo sapiens
US-09-925-548-1454

Query Match
Best Local Similarity 100.00%
Method 100.00% 100.00%

Query Match
Best Local Similarity 100.00%
Method 100.00% 100.00%

Result 10

us-09-925-548-1454/0
1 Response 145.0% Application US/09/240/6A
2 GENERAL INFORMATION:
3 APPLICANT: Compugen LTD
4 TITLE OF INVENTION: Variants of alternative splicing
5 FILE REFERENCE: 129181.4 Compugen
6 CURRENT APPLICATION NUMBER: US/09/240/6A
7 NUMBER OF SEQ ID NOS: 97222
8 SOFTWARE: Patent in version 3.2
9 SEQ ID NO: 1454
10 LENGTH: 964
11 TYPE: DNA

Query Match
Best Local Similarity 100.00%
Method 100.00% 100.00%

Query Match
Best Local Similarity 100.00%
Method 100.00% 100.00%

Result 10

us-09-925-548-1454/0
1 Response 145.0% Application US/09/240/6A
2 GENERAL INFORMATION:
3 APPLICANT: Compugen LTD
4 TITLE OF INVENTION: Variants of alternative splicing
5 FILE REFERENCE: 129181.4 Compugen
6 CURRENT APPLICATION NUMBER: US/09/240/6A
7 NUMBER OF SEQ ID NOS: 97222
8 SOFTWARE: Patent in version 3.2
9 SEQ ID NO: 1454
10 LENGTH: 964
11 TYPE: DNA
12 ORGANISM: Homo sapiens
US-09-925-548-1454

Query Match
Best Local Similarity 100.00%
Method 100.00% 100.00%

Query Match
Best Local Similarity 100.00%
Method 100.00% 100.00%

Query Match
Best Local Similarity 100.00%
Method 100.00% 100.00%